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Cancer Diagnosis and Therapy
Technical Field
The invention relates to a novel oncofetal
glycoprotein which is expressed in certain tumours
antibodies to the protein, and uses of the
antibodies in cancer diagnosis.
Background Art
The cancer phenotype typically displays loss of
differentiation, loss of proliferative control and
altered cell adhesion molecule expression. Cell
surface proteins have been shown to play an
important role in cell-cell interactions (eg NCAM),
cell-extra-cellular interactions (eg CD44) and cell
regulation (eg Notch signaling).
Some of these cell surface proteins have oncofetal
expression profiles and as such have been used as

- 1 tumour specific diagnostic markers (eg CEA). A
- 2 further use for antibodies specific for cell
- 3 surface proteins over expressed in cancer has been
- 4 in the treatment of cancer by
- 5 immunotherapy/radioimmunotherapy (eg Herceptin an
- 6 antibody recognizing HER2).

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- 8 Statements of Invention
- 9 In one aspect, the invention relates to an isolated
- 10 nucleic acid sequence which comprises a sequence
- 11 selected from the group consisting of: Sequence ID
- 12 No.1, Sequence ID No. 2, and Sequence ID No. 3.
- 13 Typically, the nucleic acid sequence is a DNA
- 14 sequence. In one embodiment, the isolated nucleic
- 15 acid sequence consists of a sequence selected from
- 16 the group consisting of: Sequence ID No. 1, Sequence
- 17 ID No. 2 and Sequence ID No. 3.

18

- 19 The invention also relates to an isolated protein
- 20 encoded by the isolated nucleic acid sequences of
- 21 the invention, or a fragment or analogue therof.
- 22 Typically, the protein is a cell surface
- 23 glycoprotein. In one preferred embodiment, the
- 24 isolated protein is an oncofetal protein expressed
- 25 by an astrocytoma cell. Typically, the protein has a
- 26 molecular weight of approximately 200kda. In this
- 27 specification, the term "protein" should be
- 28 understood as including amino acid sequences which
- 29 would more generally be referred to a peptides.

- 31 In another aspect, the invention relates to an
- 32 antibody which binds specifically to the protein of

1 the invention and any other antibody that competes directly or by stearic hindrance therewith for said 2 protein. Typically, the antibody is a monoclonal 3 antibody. In one embodiment, the antibody is a class 4 M immunoglobulin with a kappa-light chain. 5 6 7 In another aspect, the invention relates to a fragment of the antibody of the invention, which 8 fragment binds specifically to the protein of the 9 10 invention. 11 In another aspect, the invention relates to a method 12 13 of producing an antibody to a protein comprising: 14 15 innoculating an animal with a protein according 16 to the invention, wherein the protein elicits an immune response in the animal to produce the 17 18 antibody; and 19 20 isolating the antibody from the animal. 21 In one embodiment, the animal is innoculated with G-22 23 CCM cells of ECACC deposit No. 86022702. 24 In a further aspect, the invention relates to a 25 26 process for producing a hybridoma, comprising the step of innoculating a suitable subject with a 27 protein according to the invention, or an antigenic 28 29 fragment thereof, and fusing cells from the subject with a myeloma cell to produce the hybridoma. 30

Typically, the subject is innoculated with G-CCM

cells of ECACC deposit No. 86022702.

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1 In a further aspect, the invention relates to a 2 hybridoma cell obtainable according to the above 3 process. In one embodiment, the invention relates to 4 a hybridoma cell of, or derived from, ECACC Deposit 5 No. 03073001. 6 7 A deposit of hybridoma cells according to the 8 invention was made at the European Collection of 9 Cell Cultures on 30 July 2003 and accorded the 10 accession number ECACC 03073001. 11 12 In another aspect, the invention relates to a 13 monoclonal antibody obtainable from a hybridoma cell 14 of, or derived from, ECACC Deposit No. 03073001. 15 16 The invention also relates to a method of detecting 17 an astrocytoma cell in a sample of human cells, 18 which method comprises the step of contacting the 19 cell sample with an antibody of the invention, or a 20 fragment thereof, and detecting those cells which 21 have bound the antibody or fragment, wherein binding 22 of the antibody or the fragment to a cell is 23 indicative of an astrocytoma cell. Typically, the 24 antibody is a monoclonal antibody of the invention. 25 26 The invention also relates to a method of detecting 27 a primary breast carcinoma cell in a sample of human 28 cells, which method comprises the step of contacting 29 the cell sample with an antibody of the invention, 30 or a fragment thereof, and detecting those cells 31 which have bound the antibody or fragment, wherein 32

1 binding of the antibody or the fragment to a cell is

- 2 indicative of a primary breast carcinoma cell.
- 3 Typically, the antibody is a monoclonal antibody of

4 the invention.

5

- 6 The invention also relates to a diagnostic kit for
- 7 diagnosing the presence of a cell selected from the
- 8 group consisting of: astrocytoma cells; malignant
- 9 melanoma secondary tumour cells; and primary breast
- 10 carcinoma cells, the kit comprising an antibody
- 11 according to the invention, or a fragment thereof.
- 12 Typically, the antibody is a monoclonal antibody of
- 13 the invention. In one embodiment, the antibody of
- 14 the invention comprises a detectable label.
- 15 Alternatively, the kit comprises a secondary
- 16 antibody which specifically binds the antibody of
- 17 the invention, which secondary antibody comprises a
- 18 detectable label.

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- 20 The invention also relates to a biological targeting
- 21 device comprising an antibody, typically a
- 22 monoclonal antibody, of the invention, or a fragment
- 23 thereof, and a therapuetic ligand.

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- 25 The invention also relates to a therapeutic antibody
- 26 comprising an antibody, typically a monoclonal
- 27 antibody, of the invention, or a fragment thereof.

- 29 The invention also relates to a method of treating
- 30 cancer in an individual by inducing apoptosis in
- 31 cells in the individual which express a protein of
- 32 the invention, which method comprises a step of

treating an individual with an antibody of the 1 2 invention, or a fragment thereof. Typically, the antibody is a monoclonal antibody. 3 embodiment, the cancer is selected from the group 4 5 consisting of: malignant astrocytomas; malignant melanoma secondary tumours; and primary breast 6 carcinomas. Typically, the antibody is humanised. 7 8 9 The invention also relates to a polynucleotide which 10 is anti-sense to at least a portion of an insolated 11 nucleic acid sequence of the invention. Typically, 12 the polynucleotide is anti-sense to all or part of a 13 transcription initiator site of the isolated nucleic 14 acid sequence of the invention. In one embodiment, 15 the anti-sense polynucleotide comprises, or consists of, a sequence of Sequence ID No. 4. 16 17 18. The invention also relates to a method of treating 19 cancer in an individual by inducing apoptosis in 20 cells in the individual which express a protein of 21 the invention, which method comprises a step of 22 treating an individual with an anti-sense 23 polynucleotide of the invention. In one embodiment, 24 the cancer is selected from the group consisting of: 25 malignant astrocytomas; malignant melanoma secondary 26 tumours; and primary breast carcinomas. Methods of 27 delivery of anti-sense polynucleotides will be well-28 known to those skilled in the art of gene therapy. 29

30 The monoclonal antibodies of the invention may be

31 the complete antibodies described herein, or

32 fragments thereof. That is, they may be any fragment 1 of a monoclonal antibody of the invention that

- 2 specifically recognises the protein of the
- 3 invention. Such fragments include Fab, F(ab')2,
- 4 Fab', etc. These fragments ban be prepared by
- 5 digestion with an enzyme such as papain, pepson,
- 6 ficin, or the like. The properties of the obtained
- 7 fragments can be confirmed in the same manner as
- 8 described herein.

- 10 The principle reason for the poor prognosis
- 11 associated with malignant astrocytomas is recurrence
- 12 due to invasion of surrounding brain parenchyma by
- 13 tumour cells with an invasive phenotype. This
- 14 phenotype displays loss of differentiation,
- 15 secretion of proteases and altered cell adhesion
- 16 molecule expression. As part of an investigation
- 17 into the mechanisms of astrocytoma invasion,
- 18 monoclonal antibodies (Mab) were raised against cell
- 19 surface proteins expressed by an anaplastic
- 20 astrocytoma cell line (G-CCM). One of the
- 21 antibodies produced (MQ1 Mab) recognizes a
- 22 previously undescribed cell surface glycoprotein
- 23 (MQ1). In vitro MQ1 protein expression was found on
- 24 astrocytomas and fetal astrocytes, with the level of
- 25 expression increasing with astrocytoma malignancy
- 26 and decreasing with fetal astrocyte maturity.
- 27 Immunohistochemistry on histologically normal and
- 28 neoplastic brain tissue demonstrated that MQ1
- 29 protein expression is restricted to astrocytomas
- 30 (n=52). Other primary brain tumours tested
- 31 (oligodendrogliomas, neurinomas, PNET, and
- 32 medulloblastomas) and normal brain cells, including

1 neurons, oligodendrocytes and endothelial cells were

- 2 MQ1 negative, thus indicating that the MQ1 proteins
- 3 have the expression pattern of oncofetal proteins.
- 4 Similarily a study looking at primary breast
- 5 carcinomas found 60% were MQ1 positive (n=228).
- 6 Surrounding normal tissue, fibrocystic disease and
- 7 fibroadenoma tissue were MQ1 negative. Malignant
- 8 melanoma secondary tumours to the brain were also
- 9 found to be strongly MQ1 positive.

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- 11 A cDNA expression library was synthesized from G-CCM
- 12 mRNA and screened with the MQ1 antibody. Two
- 13 positive clones were isolated (Sequence ID No.s 1
- 14 and 2) and sequencing data demonstrated that both
- 15 have a high degree of homology with Jaggedl, a human
- 16 Notch ligand which plays a role in differentiation
- 17 and determination of cell fate. The library was
- 18 rescreened with probes generated from the positive
- 19 clones and further homologous transcripts were
- 20 isolated including a possible Jagged1 splice variant
- 21 (Sequence ID No. 3). Northern blotting for a range
- 22 of cell lines with these probes revealed the
- 23 presence of two transcripts (approximately 3.5kb &
- 24 5.0kb). Subsequent protein studies
- 25 (immunocytochemistry, immunoblotting and co-
- 26 immunoprecipitation) indicate that the MQ1 protein
- 27 has a high degree of homology with, but is not
- 28 identical to, Jagged1.

- 30 This investigation has identified a novel oncofetal
- 31 glycoprotein with homology to Jagged1. Its tumour
- 32 specificity together with its potential role in

regulating cellular differentiation /apoptosis 1 suggest that it may be a valuable prognostic marker and therapeutic target. 3 5 The invention will be more clearly understood from the following description of some embodiments 7 thereof, given by way of example only, with reference to the following Figures in which: 8 9 10 Fig.1A illustrates confocal microscopy of live G-CCM cells immunolabelled with MQ1 showing recognition of 11 a cell surface epitope; 12 13 Fig. 1B illustrates confocal microscopy of 14 permeabilized G-CCM cells immunolabelled with MQ1 15 16 showing recognition of an intracellular epitope and localisation of the antigen at areas of cell contact 17 18 on the cell surface; 19 Fig 2 shows a comparision of MQ1 expression, by 20 immunocytochemistry and flow cytometry, on a range 21 of fetal astrocyte cultures and astrocytoma cell 22 lines. A-C show immunocytochemistry on live cells of 23 24 a grade IV, grade III and 16 week gestation fetal astrocytes respectively. D-F chow the corresponding 25 flow analysis with the same cells with the level of 26 MQ1 surface expression estimated as mean channel 27 fluorescence. G shows the results of the flow 28 29 analysis plotted as a graph. This demonstrates an inverse correlation of cell surface MQ1 protein 30 expression with fetal astrocyte maturity and 31 32 correlation with astrocytoma grade;

1 2 Fig. 3 shows immunohistochemistry displaying diffuse MQ1 positivity throughout A) Grade I astrocytomas B) 3 Grade II astrocytomas C) Grade III astrocytomas & D) 4 focal positivity in grade IV astrocytoma cells 5 6 palisading an area of necrosis; 7 8 Fig. 4 shows MQ1 immunocytochemistry showing A) strong MQ1 positivity at the tumour front B) strong 9 MQ1 positivity in reactive astrocytes in adjacent 10 tissue C) GFAP positivity in reactive MS tissue D) 11 12 MQ1 negative reactive MS tissue; 13 14 Fig. 5 shows MQ1 immunohistochemistry of breast 15 carcinoma tissue showing A) strong MQ1 positivity in invasive ductal carcinoma cells surrounded by MQ1 16 negative stroma B) strong MQ1 positivity in lobular 17 18 carcinoma surrounded by MQ1 negative stroma; 19 Fig.6 shows MQ1 immunocytochemistry of G-CCM cells 20 21 treated with (A) 0.1µm control oligo (B) 0.5µm control oligo (C) 1.0 µm control oligo (D) 0.1 µm 22 anti-sense MQ1 oligo (E)0.5µm anti-sense MQ1 oligo 23 24 and (F) 1.0 µm antisense oligo, showing that MQ1 anti-sense oligo knocks out MQ1 protein expression 25 at concentrations of 0.5 and 1.0 µm; 26 27 Fig. 7 shows an immunoblot indicating Parp cleavage 28 of oligo-treated G-CCM cells; 29 30 31 Fig. 8 shows immunocytochemistry (ICC) detection of cleaved Caspase 3 following oligo treatmentl; and 32

1 2 Fig. 9 shows G-CCM cells labelled with MQ1 antibody 3 by ICC, 24 hours post-treatment with control and anti-sense oligonucleotides in which: 4 5 6 (A) control oligo 0.1µM 7 (B) control oligo 0.5µM (C) control oligo 1.0μM 8 9 (D) Anti-sense oligo 0.1µM (E) Anti-sense oligo 0.5μM 10 (F) Anti-sense oligo 0.1µM 11 12 13. MATERIALS AND METHODS 14 15 Materials 16 17 All cell culture reagents were obtained from Gibco 18 BRL (Paisley, UK) with the exception of the 19 hypoxanthine, aminopterin and thymidine (HAT) and the hypoxanthine and thymidine (HT) that were 20 21 obtained from Sigma (Poole, Dorset, UK). The secondary and negative control antibodies were 22 supplied by Dako (Bucks, UK). The PARP and Caspase3 23 24 antibodies were purchased from Sigma (Poole, Dorset, 25 UK) and the Protein-A Sepharose CL4B from Pharmacia Biotech (Herts, UK). PTO linked oligonucleotides 26 27 were obtained from MWG-Biotech (Germany). 28 29 Cell culture 30 31 The CB109 cell line was established from a 32 glioblastoma multiforme [6] and was a gift from Dr

- 1 Claude Chauzy (Centre Henri Becquerel, Rouen,
- 2 France). The G-CCM cell line was derived from a
- 3 human anaplastic astrocytoma and was a gift from Dr
- 4 Ian Freshney (Department of Clinical Oncology,
- 5 University of Glasgow, UK). The G-CCM cell line is
- 6 commercially available from the European Collection
- 7 of Cell Cultures under Deposit No 86022702. The
- 8 fetal astrocyte cell cultures were a gift from Ms
- 9 Kim Martin (Department of Neuropathology, Institute
- 10 of Psychiatry, London, UK). The C6 cell line,
- 11 derived from a rat glioma , was obtained from Flow
- 12 Laboratories (Scotland, UK). The skin fibroblast
- 13 cell culture was initiated in our laboratory from a
- 14 surgical specimen obtained from the Neurological
- 15 Unit (Royal Victoria Hospital, Belfast, UK). The
- 16 remaining glioma cell lines were initiated in our
- 17 laboratory from surgical specimens received from the
- 18 Neurosurgical Unit (Royal Victoria Hospital,
- 19 Belfast, UK) and were used experimentally after 5-10
- 20 passages. Tumour grading follows the World Health
- 21 Organisation classification. Cell lines were
- 22 incubated at 37oC/5% CO2 in Dulbecco's modified
- 23 Eagle's medium (DMEM) containing 2mM glutamine, 10%
- 24 fetal calf serum (FCS), and phenol red. All cell
- 25 lines were tested for mycoplasma using Hoechst 33258
- 26 fluorescent dye and were found to be negative.

27

28 Monoclonal antibody production

- 30 Mabs were produced utilizing a standardized protocol
- 31 designed to promote a rapid predominantly IqG
- 32 response. In brief, a BALB/c mouse was inoculated

1 intra-peritoneally with 5x106 G-CCM cells in 1ml of

- 2 Freund's complete adjuvant. Similar doses
- 3 emulsified in Freund's incomplete adjuvant were
- 4 administrated 14 and 28 days later to boost the
- 5 immune response. Four days after the final booster
- 6 inoculation the mouse was killed, its spleen
- 7 aseptically removed and the splenocytes induced to
- 8 fuse with NSO myeloma cells (at a ratio 5:1) using
- 9 polyethylene glycol. The resulting fusion products
- 10 were suspended in a selective, HAT-supplemented,
- 11 growth medium (RPMI-1640 medium containing 10mM L-
- 12 glutamine, 1% sodium pyruvate, 100 iu/ml penicillin,
- 13 100□g/ml streptomycin and 20% Myoclone FCS) and
- 14 seeded into 96-well plates. The medium, from the
- 15 viable hybridomas produced, was screened by indirect
- 16 immunofluorescence against live and acetone-fixed G-
- 17 CCM cells. Those showing specific recognition were
- 18 recloned three times, to ensure monospecificity, in
- 19 HT-supplemented growth medium and stored in liquid
- 20 nitrogen. The hybridoma cell line MQ-1, which
- 21 produced an antibody recognizing a cell surface
- 22 antigen was propagated as an ascitic tumour in
- 23 BALB/c mice previously immunosuppressed with
- 24 Pristane. The ascitic fluids were collected,
- 25 centrifuged and frozen at -20oC until use.

26

- 27 The positively labelling Mabs were isotyped for
- 28 their class and light chains using a monoclonal
- 29 antibody isotyping kit.

30

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1 Immunofluorescence 2 Hybridoma medium (neat) or ascites fluid (diluted 3 1:200 in PBS) was incubated with living cells, grown 4 5 to 90% confluence on coverslips, for 40 min at room 6 temperature (RT). After washing, the cells were fixed in acetone at -20oC for 10 min followed by 7 rehydration in PBS and incubation with an FITC-8 9 conjugated rabbit antimouse antibody (FITC-RAM) for 10 30 min at RT. After two further washes the cells were mounted on a glass slide, in a drop of 11 Citifluor, and examined using a Zeiss 12 13 immunofluorescence microscope or a Biorad confocal 14 microscope. Incubations in PBS without primary 15 antibody were used as negative controls. 16 fluorescent labelling of positive cells was 17 subjectively rated from low intensity (+) to high 18 intensity (++++). 19 20 Flow Cytometry 21 22 A preliminary study (results not shown) comparing 23 the expression of MQ-1 protein on cells removed 24 enzymatically (trypsin) and non-enzymatically (0.53 25 mM EDTA in PBS) from culture flasks, revealed that 26 the MQ1 protein epitope was trypsin-resistant. 27 28 Cultured cells were removed from the flasks by 29 trypsinization, counted and aliquoted into 30 centrifuge tubes at a concentration of 5x105 cells

Triplicate samples were incubated in

excess ascitic fluid in 2000l of serum free medium

1 containing 1% bovine serum albumin (SFM/BSA) for 40

- 2 min at RT with gentle agitation. Following 2
- 3 washes in SFM the cells were incubated in an FITC-
- 4 RAM antibody for 30 min at RT with gentle
- 5 agitation. The cells were then washed twice in SFM
- 6 and fixed in PBS containing 1% para-formaldehyde.
- 7 The samples were analysed within 48 hr of fixation,
- 8 using a Coulter EPICS Elite flow cytometer.
- 9 Negative controls were incubated with an antibody
- 10 raised against Aspergillus niger glucose oxidase, an
- 11 enzyme not present or inducible in mammalian cells.
- 12 The consistency of the mean channel fluorescence
- 13 measurements between sample batches was checked
- 14 using EPICS Immuno-Brite standards.

15

16 Immunohistochemistry

- 18 On receipt the tissue was fixed in 10% formalin
- 19 prior to routine embedding in paraffin wax using a
- 20 Tissue Tex VIP (Miles Scientific) automated
- 21 processor. The paraffin blocks were sectioned at a
- 22 thickness of 6mm and mounted onto 3-
- 23 aminopropyltriethoxysilane-coated slides. The
- 24 tissue sections for indirect immunohistochemistry
- 25 were processed using an avidin-biotin peroxidase
- 26 complex (ABC) method. The tissue was dewaxed in
- 27 xylene and rehydrated before endogenous peroxidase
- 28 activity was blocked by a 10min incubation in 3%
- 29 H2O2 in methanol at room temperature (RT). To
- 30 counter antigen masking, due to the formalin
- 31 fixation, the tissue was pretreated with microwave
- 32 irradiation to promote antigen retrieval. The

1 sections were washed in distilled water and placed

- 2 in 0.01M Tri-Na citrate pH7.8 and irradiated in a
- 3 Miele microwave oven for 6min (2x3min) at 450W (the
- 4 optimal toime and intensity of irradiation was
- 5 determined from preliminary studies). After
- 6 incubation in PBS containing 5% normal rabbit serum
- 7 for 10min at RT the sections were incubated in MQ1
- 8 ascites (diluted 1:50 in PBS) at 4C overnight.
- 9 Following 2x5min washes in PBS the sections were
- 10 incubated in biotinylated rabbit anti-mouse IgM
- 11 diluted 1:400 in PBS for 40min at RT. After further
- 12 washes in PBS, a streptavidin-biotin complex linked
- 13 to peroxidase was added to the sections and
- 14 incubated for 40min at RT. The peroxidase reaction
- 15 was developed in 0.1% diaminobenzidine in PBS
- 16 activated with 1% H2O2. After washing in water, the
- 17 sections were counterstained in haematoxylin,
- 18 dehydrated through graded alcohols, cleared in
- 19 xylene and mounted in DPX. In addition to negative
- 20 controls, incubated with a primary antibody raised
- 21 against Aspergillus niger glucose oxidase, positive
- 22 controls of histologically normal brain and
- 23 astrocytoma tissue were included with every batch.
- 24 cDNA Expression Library and screening.

25

26 G-CCM Cell cDNA Library Synthesis

27

- 28 A Total RNA isolation from G CCM cells
- 29 This was performed using Tel-Test RNA Stat-60,
- 30 following their guidelines. Web Site
- 31 www.isotexdiagnostics.com/rna stat-60 reagent.html

1	B mRNA Purification from Total RNA
2	This was performed using Invitrogen's FastTrack
3	2.0 Kit, following their guidelines. Web Site
4	www.invitrogen.com/content.cfm?pageid=3443&cfid=3308
5	35&cftoken=53475959#FastTrack
6	
7	C cDNA Library Synthesis from mRNA
8	This was performed using a Stratagene cDNA
9	synthesis kit (following their protocol).
10	Stratagene ZAP Express cDNA Synthesis Kit
11	Instruction Manual
12	www.stratagene.com/manuals/200403.pdf
13	·
14	RESULTS
15	
16	Antibody Production
17	
18	The fusion resulted in the production of five viable
19	antibody secreting hybridomas which screened
20	positively by immunofluorescence microscopy on
21	acetone fixed G-CCM cells. Of these, one (hybridoma
22	MQ1) was found to secrete an antibody which was
23	isotyped as a class M immunoglobulin with a kappa-
24	light chain. This antibody recognizes a cell
25	surface epitope, showing punctate labelling, on live
26	G-CCM cells. Further examination by confocal
27	microscopy confirmed the cell surface labelling of
28	live G-CCM cells and revealed the presence of an
29	intra-cellular epitope in permeabilized cells
30	(Figure 1A&B). In addition examination of the
31	permeabilized cells demonstrated localisation of
32	labelling at focal adhesion points on the cell

surface. 1

2

Immunocytochemistry 3

4

- A range of cell lines was examined by indirect 5
- immunofluorescence for the presence of the MQ-16
- antigen 7
- (Table 1). 8

9

Table 1

CELL LINE	TISSUE SOURCE	MQ1	LABELLING
Fibroblasts	Normal skin		
•			_
C6	Rat glioma		
FA 10 weeks	Human fetal astrocytes		+
FA 12 weeks	Human fetal astrocytes		+
FA 14 weeks	Human fetal astocytes		+
FA 15 weeks	Human fetal astrocytes		+
FA 16 weeks	Human fetal astrocytes		+
FA 19 weeks	Human fetal astrocytes		+
NP 527/94	Pilocytic astrocytoma (I)		++
NP 396/94	Pilocytic astrocytoma (I)		++
NP 424/94	Astrocytoma (II)		++
NP 676/92	Astrocytoma (II)		++
NP 445/92	Astrocytoma (II)		++
NP 204/92	Astrocytoma (II)		++
NP 482/96	Astrocytoma (II)		++
NP 473/92	Anaplastic astrocytoma (III)		+++
G-CCM	Anaplastic astrocytoma (III)		++++
NP 493/94	Anaplastic astrocytoma (III)		+++
NP 785/96	Anaplastic astrocytoma (III)		+++

++++	(IV)	${\tt multiforme}$	Glioblastoma	402/93	NP
. +++	(IV)	${\tt multiforme}$	Glioblastoma	293/96	NP
++++	(IV)	${\tt multiforme}$	Glioblastoma	602/91	NP
+++	(VI)	${\tt multiforme}$	Glioblastoma	536/94	NP
++++	(VI)	${\tt multiforme}$	Glioblastoma	306/92	NP
+ + +	(VI)	${\tt multiforme}$	Glioblastoma	479/95	NP
+ + +	(IV)	${\tt multiforme}$	Glioblastoma	770/96	NP
+ + + +	(VI)	${\tt multiforme}$	Glioblastoma	876/96	NP
+ + +	(VI)	${\tt multiforme}$	Glioblastoma	39/96	NP
_	(VI)	${\tt multiforme}$	Glioblastoma	109	СВ
_	(IV)	multiforme	Glioblastoma	670/92	NP

Table 1. Indirect immunofluorescence on a range of live cell lines and cell cultures with MQ1 antibody. 2 3 The results show that the human skin fibroblasts and 4 the C6, rat glioma, cell lines do not express the 5 antigen. The fetal astrocytes and glioma cell lines 6 were positive with the exception of two cell lines 7 (CB109 and NP670/92) derived from glioblastomas 8 multiforme. Under subjective microscopic analysis 9 there appeared to be a variation in labelling 10 intensity between the positive cell lines. The high 11 grade gliomas had a higher labelling intensity than 12 This was low grade gliomas and fetal astrocytes. 13 confirmed by flow cytometry (Figure 2). 14 show a progressive increase in MQ-1 antigen 15 expression, as estimated by the mean channel 16 fluorescence, from low to high grade astrocytomas, 17 the expression on grade IV astrocytomas being more 18 than double that of grade I astrocytomas. The fetal 19 astrocytes showed a lower expression than the 20 astrocytoma cell lines, that halved from fetal 21 astrocytes of 12 weeks gestation to 16 weeks 22 gestation. 23 24 Immunohistochemistry 25 26

29

The results of the immunohistochemical study on 27 primary brain tumours are summarized in Table 2. 28

Table 2

Tumour	# Biopsies	MQ1 positivity
Astrocytomas	30	29/30
Neurinoma	3	0/3
Oligodendroglioma	, 3	0/3
Medulloblastoma	3	0/3
PNET	3	0/3

- 1 Table 2 Immunohistochemical analysis of MQ1 immuno-
- 2 labelling of a range of Primary Brain Tumours
- 3 showing that of the tumour tissue tested only
- 4 astrocytomas displayed MQ1 positivity.

- 6 The results show that of all the primary brain
- 7 tumours tested (oligodenrogliomas, PNET etc) only
- 8 astrocytomas were MQ1 positive.
- 9 All pilocytic (grade I) astrocytomas showed a
- 10 similar staining pattern. There was strong cellular
- 11 immunostaing of MQ1 proteins which extended to the
- 12 cellular processes of bipolar cells (Fig3A). The
- 13 immunopositive cells stood out prominently against a
- 14 loosely arranged less cellular stroma.
- 15 The astrocytomas (grade II) and anaplastic (grade
- 16 III) astrocytomas revealed a diffuse
- 17 immunopositivity and the staining pattern was
- 18 similar in all (Fig 3B&C). There was variation in
- 19 the staining pattern of glioblastomas. Out of 16
- 20 glioblastomas tested, 1 was unreactive revealing no
- 21 MQ1 protein expression whereas 14 showed focal
- 22 positivity and one diffuse immunostaining (Fig 3D).
- 23 Focal positivity was observed as clusters or groups

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- of positive cells surrounded by unreactive aresas. 1
- 2 Tumour cells palisading around areas of necrosis, a
- characteristic feature of glioblastomas also reveled 3
- focal positivity. However tumour giant cells, 4
- bizarre cells and clusters of proliferating
- 6 endothelial cells were negative for MQ1 protein
- expression. The oligodendroglial cells were 7
- 8 negative. Within adjacent grey matter the neurones
- did not show immunolabelling for the MQ1 proteins. 9
- 10 The endothelial cells lining small and large blood
- 11 vessels in and around tumours of all grades showed
- 12 no MQ1 protein expression. There was no
- immunolabelling of lymphocytes in the perivascular 13
- The infiltrating edge of the tumours and 14
- 15 the adjacent glial areas showed prominent labelling
- of large reactive astrocytes (Fig 4 A&B)). 16
- cells revealed multiple processes. However this MQ1 17
- positivity in reactive astrocytes was only found 18
- surrounding MQ1 positive tumours, other reactive 19
- 20 tissue such as MS tissue that shows prominent
- reactive astrocytes when labeled for GFAP (FIG 4C) 21
- displayed no MQ1 positivity in the 10 biopsies 22
- 23 tested (Fig 4D).
- In non-CNS tissue tested malignant melanoma and 24
- 25 breast 20 to the brain were found to express the MQ1
- proteins (Table 3). 26

Table 3

Tissue	# Biopsies	MQ1 Positivity
Breast 20 (brain)	3	3/3
Breast 10	228	137/228

	Fibroadenoma 5	0/5
	Fibrocystic Diease 5	0/5
	M.Melanoma2o (brain) 4	4/4
1		
2	Table 3 Immunohistochemical MQ1 immunolabelling of a	
3	range of non-CNS tumours, showing MQ1 positivity in	
4	60% of primary breast tumours and no positivity in	
5	fibrocystic diease and fibroadenomas that are non-	
6	malignant breast conditions.	
7		
8	Of the primary breast tumours tested 137/228 were	
9	MQ1 positive while fibrocystic diease and	
10	fibroadenoma tissues, both premalignant conditions	
11	displayed no MQ1 positivity. Figure5 shows strong	
12	MQ1 positivity in invasive ductal carcinoma cells	
13	and lobular carcinoma cells surrounded by MQ1	
14	negative stroma.	
15		
16	Isolation of MQ-1 Clones	
17		
	•	
18	Screening of a cDNA expression library (from G-CCM	
18 19	Screening of a cDNA expression library (from G-CCM mRNA) with the MQ1 antibody identified two clones	
19	mRNA) with the MQ1 antibody identified two clones	
19 20	mRNA) with the MQ1 antibody identified two clones with significant homology to the Jagged 1 protein	
19 20 21	mRNA) with the MQ1 antibody identified two clones with significant homology to the Jagged 1 protein	
19 20 21 22	mRNA) with the MQ1 antibody identified two clones with significant homology to the Jagged 1 protein (Sequence ID No's 1 and 2).	
19 20 21 22 23	mRNA) with the MQ1 antibody identified two clones with significant homology to the Jagged 1 protein (Sequence ID No's 1 and 2).	
19 20 21 22 23 24	mRNA) with the MQ1 antibody identified two clones with significant homology to the Jagged 1 protein (Sequence ID No's 1 and 2). Antisense Treatment Protocol	
19 20 21 22 23 24 25	mRNA) with the MQ1 antibody identified two clones with significant homology to the Jagged 1 protein (Sequence ID No's 1 and 2). Antisense Treatment Protocol Antisense Oligonucleotide	
19 20 21 22 23 24 25 26	mRNA) with the MQ1 antibody identified two clones with significant homology to the Jagged 1 protein (Sequence ID No's 1 and 2). Antisense Treatment Protocol Antisense Oligonucleotide	

- 2 The PTO linked antisense oligonucleotide was
- designed against the transcription initiation site 3
- and kozac sequence at the beginning of the Jagged1 4
- gene (Accession number AF028593). The control 5
- oilgonucleotide was the same 18 mer with one base 6
- changed (therefore being the tightest control 7
- possible to generate). Both oligonucleotides were 8
- synthesized by MWG Biotech. For colony count assays 9
- 10 G-CCM cells were seeded out into 24well plates at
- 50,000 cells/well. The cells were incubated for 11
- 24hrs in growth medium and then washed with serum 12
- free medium (SFM). The cells were then either 13
- treated with lipofectin (Invitrogen Life 14
- Technologies) alone following the standard protocol 15
- (at 5µl/ml) or lipofectin with the antisense and 16
- antisense control oligonucleotides at a range of 17
- 18 concentrations (0.1, 0.5 and 1.0 µM) for 16hrs.
- Following treatment the cells were washed twice with 19
- SFM and then incubated in growth medium for 24 and 20
- 48hrs. The results (Figure 6) show that treatment 21
- with the antisense oligonucleotide at concentrations 22
- of 0.5 and 1.0 µM reduced the tumour cell population 23
- when compared to the control oligonucleotide and 24
- lipofectin alone treatment. To assess whether this 25
- was due to the induction of apoptosis similarly 26
- treated cells were harvested for their protein and 27
- examined for Parp cleavage (an indicator of 28
- 29 apoptosis) by immunoblotting. The results (Figure
- 7) clearly show a reduction in the level of Parp at 30
- 0.5 and 1.0µM antisense oligonucleotide treatment 31
- when compared to control oligonucleotide and 32

1 lipofectin alone treatment. Thus indicating that

- 2 the antisense oligonucleotide treatment induces
- 3 apoptosis in the G-CCM cells. To confirm this,
- 4 treated G-CCM cells were also examined for the
- 5 presence of cleaved Caspase 3 (another indicator of
- 6 apoptosis) by immunocytochemistry. The results
- 7 (Figure 8) show that G-CCM cells treated with 1.0µM
- 8 displayed caspase 3 cleavage thus indicating that
- 9 apoptosis was being induced. To demonstrate that
- 10 these effects were due to the knocking out of the
- 11 MQ1 proteins by the antisense oligonucleotides,
- 12 treated cells were examined for the presence of the
- 13 MQ1 proteins by immunocytochemistry with the MQ1
- 14 antibody. The results (Figure 9) show that the
- 15 expression levels of the MQ1 proteins is reduced by
- 16 antisense oligonucleotide treatment when compared to
- 17 the control oligonucleotide.

18

- 19 The invention described herein has potential uses as
- 20 a:

21

- 22 Diagnostic Tool- The antibody clearly distinguishes
- 23 astrocytomas from other primary brain tumours,
- 24 normal cells and reactive gliosis. In addition it
- 25 recognizes 60% of primary breast tumours tested.

26

- 27 Targeting Device- The specificity of the antibody
- 28 means it can be used as a targeting device such as
- 29 in radioimmunotherapy.

- ·31 Therapeutic Target- The antibody itself can be used
- 32 as a therapeutic agent by blocking out signaling

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- 1 through the MQ1/Notch pathway thus inducing
- 2 apoptosis in astrocytoma cells.

- 4 The invention is not limited to the embodiments
- 5 hereinbefore described which may be varied without
- 6 departing from the spirit of the invention.